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IFWO

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 292 Lys Lys His His Val Met
293 1
 296 <210> SEQ ID NO: 14
 297 <211> LENGTH: 6
 298 <212> TYPE: PRT
 299 <213> ORGANISM: Artificial sequence
                                             -Invalid Response
 301 <220> FEATURE:
 302 <223> OTHER INFORMATION: Hexapeptide
. 305 <220> FEATURE:
 306 <221> NAME/KEY: Peptide
 307 <222> LOCATION: (1)..(6)
 308 <223> OTHER INFORMATION: Hexapeptide
310 <220> FEATURE:
311 <221> NAME/KEY: Peptide
312 <222> LOCATION: (1)..(6)
 313 <223> OTHER INFORMATION: P3
315 <400> SEQUENCE: 14
317 Lys Lys Ris Tyr Lys Met
318 1
321 <210> SEQ ID NO: 15
322 <211> LENGTH: 6
323 <212> TYPE: PRT
324 <213> ORGANISM: Artificial sequence
                                                -Invalid Response
326 <220> FEATURE:
327 <223> OTHER INFORMATION Hexapeptide
330 <220> FEATURE:
331 <221> NAME/KEY: Peptide
332 <222> LOCATION: (1)..(6)
333 <223> OTHER INFORMATION: P4
335 <400> SEQUENCE: 15
337 Lys Lys His Tyr Pro Met
338 1
341 <210> SEQ ID NO: 16
342 <211> LENGTH: 6
343 <212> TYPE: PRT
344 <213> ORGANISM: Artificial sequence
                                                Invalid Response
346 <220> FEATURE:
347 <223> OTHER INFORMATION: Hexapeptide
350 <220> FEATURE:
351 <221> NAME/KEY: Peptide
352 <222> LOCATION: (1)..(6)
353 <223> OTHER INFORMATION: P5
355 <400> SEQUENCE: 16
357 Lys Lys His Tyr Val Met
                                             The type of errors shown exist throughout
361 <210> SEQ ID NO: 17
                                           the Sequence Listing. Please check gubsequent
```

sequences for safety chars.

DATE: 02/18/2004

RAW SEQUENCE LISTING

RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/10/774,147

DATE: 02/18/2004 TIME: 15:00:46

Input Set: A:\10050-03U8A.ST25.txt
Output Set: N:\CRF4\02182004\J774147.raw

#### Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; Xaa Pos. 1,3,4,5 | Seq#:2; Xaa Pos. 4,5,6 | Seq#:3; Xaa Pos. 4,5,6 | Seq#:4; Xaa Pos. 4,5,6 | Seq#:5; Xaa Pos. 4,5,6 | Seq#:6; Xaa Pos. 1,2,3 | Seq#:8; Xaa Pos. 1,2,3 | Seq#:9; Xaa Pos. 1,2,3 | Seq#:10; Xaa Pos. 1,2,3 | Seq#:11; Xaa Pos. 1,2,3